

non-sense primer	3'-	ATACTCTGACTCTGGGGGAGACA	-51
sense primer	5'-	TGAGACC <u>C</u> CCTCTGTTATCCCAG	-3 '
Bet v 1 non-sense	3'-	${\tt TTAATACTCTGACTCTGG\underline{T}GGAGACAATAGGGTCGTCGAGC}$	-5 '
Bet v 1 sense	5'-	${\tt AATTATGAGACTGAGACC} \underline{{\tt A}CCTCTGTTATCCCAGCAGCTCG}$	-3 '





all	sense	1: 183Bv, 15-mer 5'-GTTGCCAACGATCAG
1	sense	2: 184Bv, 23-mer 5'-TGAGACCCCCTCTGTTATCCCAG
1	non-sense	3: 185Bv, 23-mer 5'-ACAGAGGGGGTCTCAGTCTCATA
2	sense	4:186Bv, 31-mer 5'-GATACCCTCTTTCCACAGGTTGCACCCCAAG
2	non-sense	5: 187Bv, 31-mer 5'-ACCTGTGGAAAGAGGGTATCGCCATCAAGGA
3	sense	6: 188Bv, 23-mer 5'-AACATTTCAGGAAATGGAGGCC
3	non-sense	7: 189Bv, 23-mer 5'-TTTCCTGAAATGTTTTCAACACT
4	sense	8: 190Bv, 23-mer 5'-TTAAGAACATCAGCTTTCCCGAA
4	non-sense	9: 191Bv, 23-mer 5'-AGCTGATGTTCTTAATGGTTCCA
5	sense	10: 192Bv, 23-mer 5'-GGACCATGCAAACTTCAAATACA
5	non-sense	11: 193Bv, 23-mer 5'-AGTTTGCATGGTCCACCTCATCA
6	sense	12: 194Bv, 23-mer 5'-TTTCCCTCAGGCCTCCCTTTCAA
6	non-sense	13: 195Bv, 23-mer 5'-AGGCCTGAGGGAAAGCTGATCTT
7	sense	14: 196Bv, 24-mer 5'-TGAAGGATCTGGAGGGCCTGGAAC
7	non-sense	15: 197Bv, 24-mer 5'-CCCTCCAGATCCTTCAATGTTTTC
8	sense	16: 198Bv, 24-mer 5'-GGCAACTGGTGATGGAGGATCCAT
8	non-sense	17: 199Bv, 24-mer 5'-CCATCACCAGTTGCCACTATCTTT
all	non-sense	18: 200Bv, 15-mer 5'-CATGCCATCCGTAAG

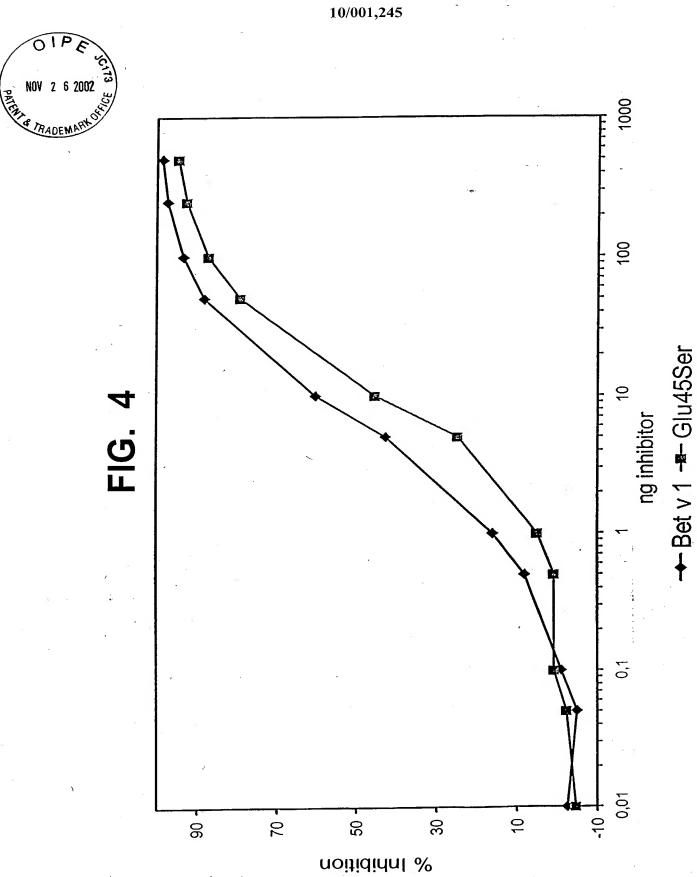


FIG. 3

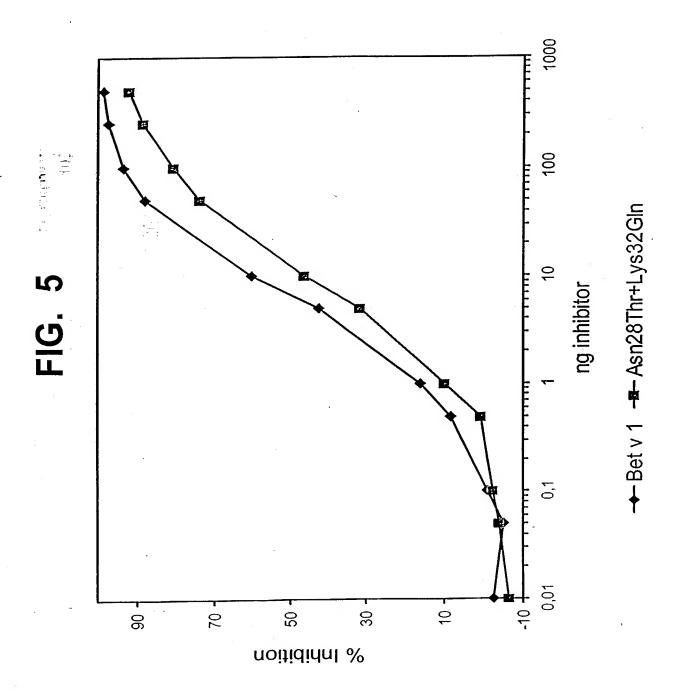
1 (A-C)

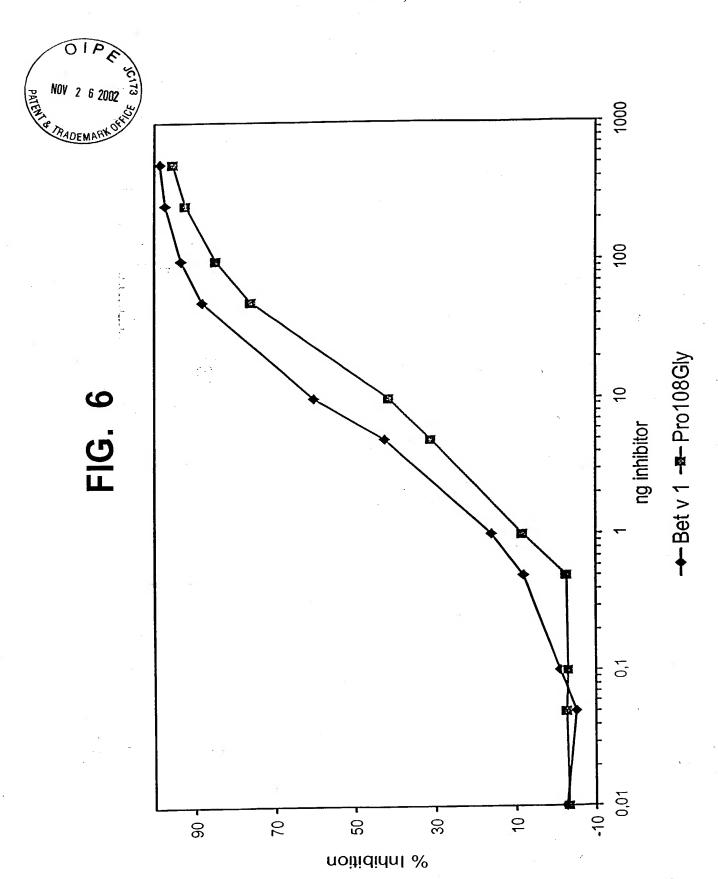
GG	TGT	GTI	TAI	ATTA	TGF	\GA(CTG	AGA(CCAC	CTC	CTG	rta:	rcc	CAG	CAG	CTC	GAC	TG	TTC	CAAG	60	
G	v	F	1	1 Y	E	9 9	r E	3 7	r T -	-P 5	, i	, :	E 1	P 2	A 2	Α. :	R	L	F	ĸ	20	
				9 (A	-G)		2	(A-	·C)	2 (A	C)											
GC	CTT	TAT	CCI	TG <u>A</u>	TGG	CGF	<u>K</u> AT/	CCI	CTI	TCC	A <u>A</u> A	GGI	TGC	ACC	:ccz	LAG(CA	TT/	AGC	AGT	120	
A	F	1	L	D-	G G		N-	T L	F	P	к-	Q V	7 A	E	. (2 #	A :	I	s	s	40	
		3 (6	GA-	TC)		7 (A	A-T	C) '					, 4	(G-	C)			6 (G	- A	TC)		
GT	'GAI	AAA	CAT	T <u>GA</u>	AGG	А <u>Д</u> А	TGG.	AGG	GCC	TGG.	AAC	CAT	TAA	GAA	<u>G</u> AT	CAG	CT	rrc	cc	<u>GA</u> A	180	
v	E	N	I	E-5	G G	N-	S G	G	₽	G	T	1	ĸ	K-	N I	s	: 1	F	P	E-S	60	
								5 (CA-TG) AGGACAGAGTTGATGAGGTGGACCA <u>CA</u> CAAACTTCAAA 240														
3GC	CTC	CC1	TT	CAAC	TAC	GTY	GAAC	GAG	CAG	\GT7	rga:	rga(GG T (GGA	CCA	<u>CA</u> C	AAA	CT	TC	AAA	240	
G				·ĸ										D		т				к	80	
'AC	AAT	TAC	AGC	GTG	ATC	GAC	GGG	GG7	ccc	ATA	GGC	GAC	CAC	YTT	GA (SAAG	GAT	CTC	CCA	AC	300	
Y	N	Y	s	v	I	E	G	G	P	I	G	D	т	L	E	K	1	8	S	N	100	
0 (gag	-CA	.C)		8	(CC	C-T	GG)														
				GTG						.GGA	TCC	ATC	TTG	AAG	ATO	AGC	CAA	CAZ	AGT	'AC	360	
E	I	ĸ	I	v	A	т	P-G	D.	, G	G	s	I.	L	ĸ	1	s	N	F	C	Y	120	
AC	ACC	AAA	GGT	GAC	CAT	GAG	GTG	AAG	GCA	GAG	CAG	GTT	'AAG	GCA	AGI	'AA'	\GA	raa	rgg	GC	420	
Н	т	ĸ	G	, D	Н	Е	v	ĸ	A	E	Q	v	к	A	s	к	Е	M	1	G.	140	
AGA	CAC	CTT	TTG.	AGG	GCC	GTT	GAG.	AGC	TAC	CTC	TTG	GCA	CAC	TCC	GAT	GCC	TAC	CAA	CT.	AA	480	
Е	T .	L	L	R	A	v	E	s	Y	L	L	A	н	s	D	A	. Y	N	8	top	159	

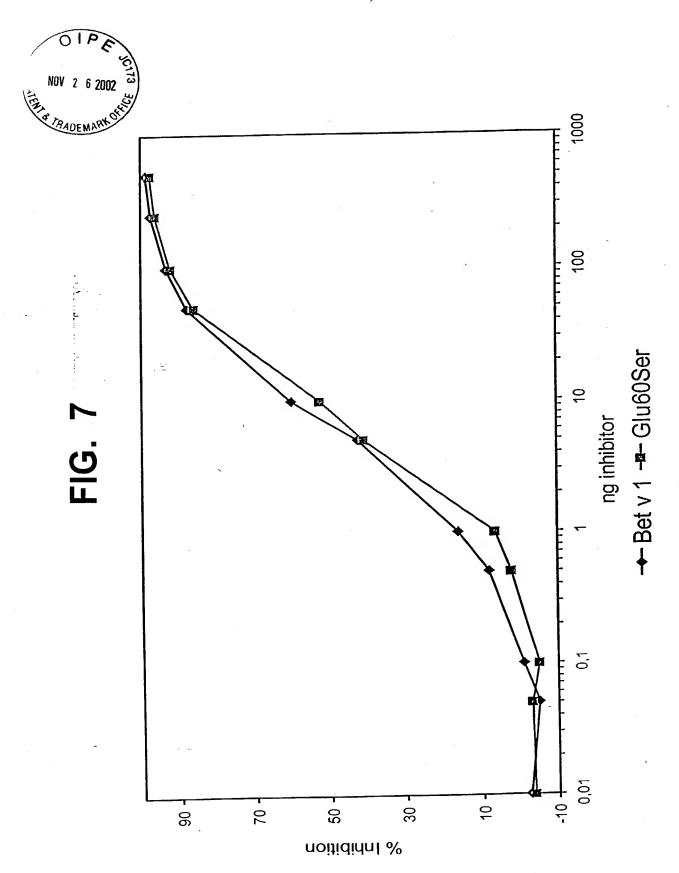




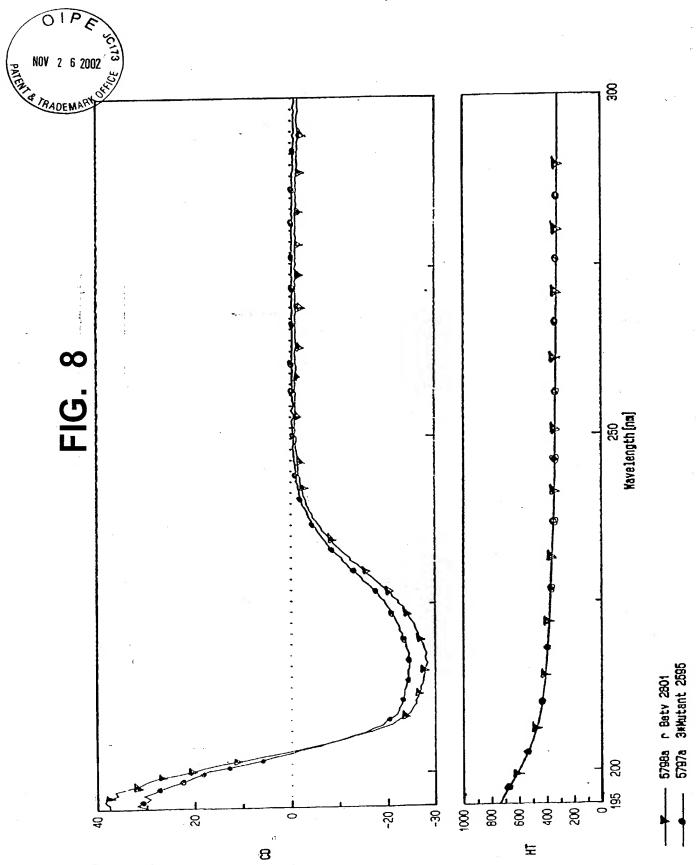












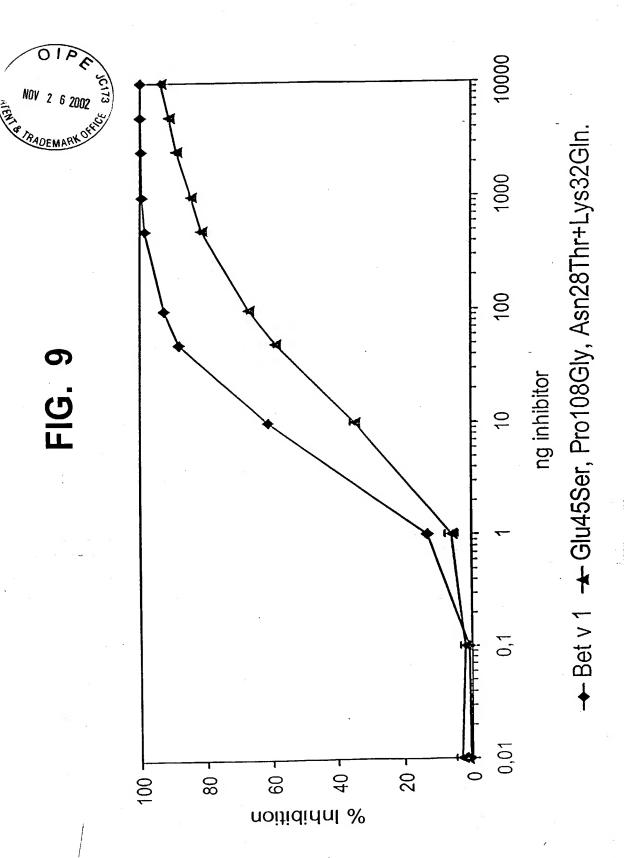




FIG. 10 A

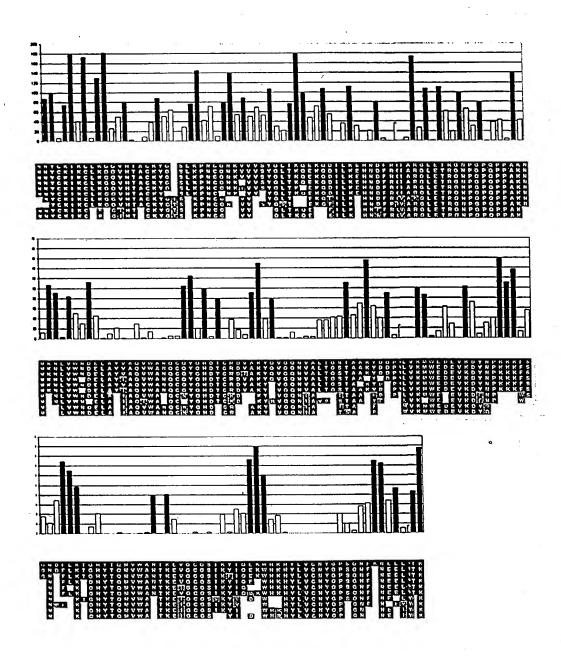




FIG. 10 B

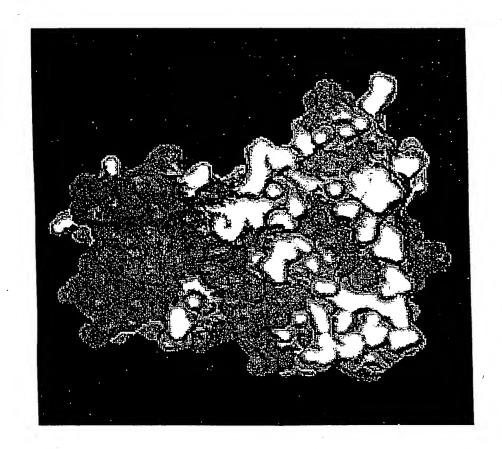




FIG. 11 A

Ves v 5 mutant 1 (K72A)

Ves v 5 sense 5'- ACCACAGCCTCCAGCGAAGAATATGAAAAATTTGGTATGGA -3'

Ves v 5 non-sense 3'- TGGTGTCGGAGGTCGCTTCTTATACTTTTTAAACCATACCT -5'

sense primer 5'- CCAGCGGCTAATATGAAAAAT -3'

non-sense primer 3'- GTCGGAGGTCGCCGATTATAC -5'

FIG. 11 B

Ves v 5 mutant 2 (Y96A)

Ves v 5 sense	5´- GG	CTAATCAATGTCAATATGGTCACGATACTTGCAGGGAT	G -3´
Ves v 5 non-sense	3'- CC	GATTAGTTACAGTTATACCAGTGCTATGAACGTCCCTA	C -5
sense primer	5 -	TGTCAAGCTGGTCACGATACT	-3 ~
non-sense primer	3 ~ -	TTAGTTACAGTT <u>CG</u> ACCAGTG	-5´

FIG. 12

all sense 1: XhoI start, 38-mer:

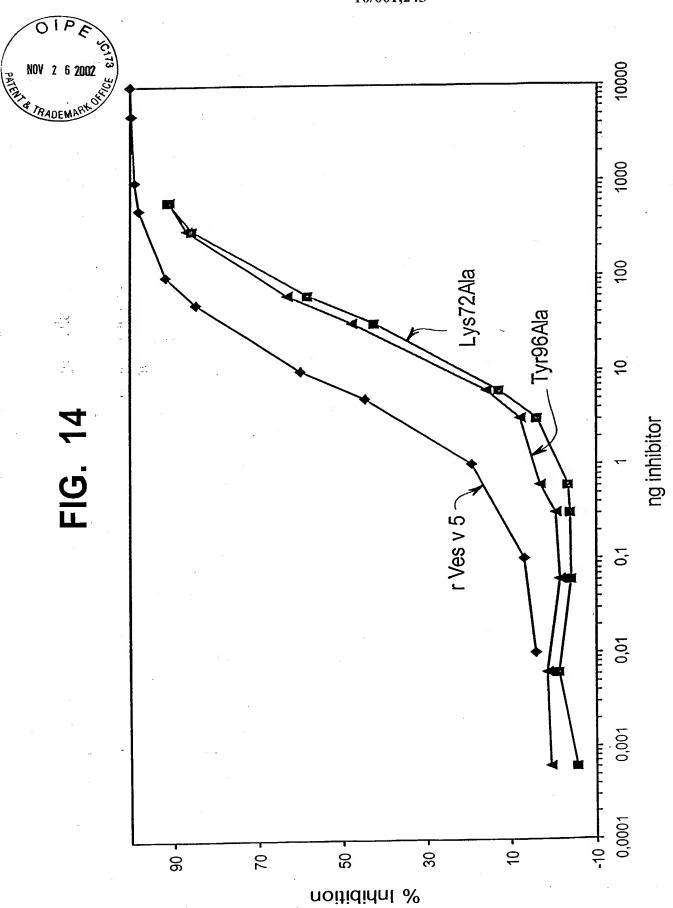
ECORI 5'-CCGCTCGAGAAAAGAAACAATTATTGTAAAATAAAATG L E K R N N Y C K I K Kex2 cleavage site amino terminus of Ves v 5

1 sense		1-mer	5´-CCAGCGGCTAATATGAAAAAT
1 non-sense		1-mer	5´-CATATTAGCCGCTGGAGGCTG
2 sense	J. 250	1-mer	5´-TGTCAAGCTGGTCACGATACT
2 non-sense		1-mer	5´-GTGACCAGCTTGACATTGATT
all non-sense	7: CT-pPICZαA	, 21-mer	5 ~ - ATTCATCAGCTGCGAGATAGG



1	AA	CAA'	TTA'	TTG	TAA	AAT	AAA	ATG:	rtt	SAA Z	AGG2	AGGT	GTC	CCAT	rac?	rgco	CTG	CAA	ATA	rgga	. 60
1	N	N	Y	C	K	I	K	C	L	K	G	G	v	H	T	A	C	K	Y	\mathbf{G}	20
			•																		
	AG	n 0 mr			333 5	DOD CI	300	חא אר	DN N/	יתייי	\ Cmc	יישי	TO	ח א יחיב	הממי	ירים	\	וממב	י אי	ימאמ	120
	AG'																				40
21	S	ь	K	Р	N	C	G	N	ĸ	V	V	V	5	1	G	ъ	1	K	Q	ь	40
101	AAA	(C) 1	CAC	ጉእ ጥ(ኅብጣን	אממי	2G N G	יכיאכ	יממי	יינאַר	adada	מאמ	ממט	מממ	רדמ	GCE	\CG!	AGGI	\ \ TTC	GAG	180
121	K		יאטני	T	- T T T	v.	ישטעני	II	וניננית א	n	F	D.	<u> </u>	ĸ	т-	A	R	G	τ.	E	60
41	K	Ų	D	1	ы	K	E	n	14	D	F	K	V	10	_	-		Ü			
														[A			-				
181	ACT	AGA	AGGT	raat	rcci	GGI	ACC A	CAC	CCI	CCA	GCG	<u>AAG</u>	PAA	ATG	AAA	raa.	TTC	GTA	TGC	BAAC	240
61	T	R	G	N	P	G	P	Q	P	P	A	K	N	M	K	N	L	V	W	N	80
																2 [¥96	[A]	TA-	GC)	
241	GAC	GAG	:ጥጥ	AGCT	TAT	YTTC	GCC	CAA	GTG	TGG	GCT	'AAT	CAA	TGT	CAA	TAT	GGI	CAC	GAT	ACT	300
81	D	E	Τ.	A	v	v	A.	0	v	W	A	N	0	C	0	Y	G	н	D	Ť	100
01		_	_		-	•		*					_	_	_						
301	TGC	'AGG	ינישטי	מידיני	יפרא	מממ	ייעריי	CAG	ĊТ	YGGA	CAA.	AAC	GTA	GCC	тта	ACA	GGI	'AGC	ACG	GCT	360
101	C	D D	ת	VÍ	Δ Δ	K	v	0	v	G.	0	N	v	A	T.	т	G	S		Α	
101	C	K	ט	·	•	10	•	×	•	. •	×		•		_	_	_				
							1		~~~					~~~	~~~	~ n n	ama	(N	~13 m	ern a ern	420
361	GCT	AAA	TAC	'GA'I	GAT	'CCA	GTT	AAA	CTA	GTT	'AAA	ATG	TGG	GAA	GAT	GAA	GIG	AAA	GAI	TAL	140
121	A	K	Y	D	D	Р	V	K	L	V	K	М	W	Е	D	В	V	K.	ע	x	140
421	TAA	ÇCT	'AAG	AAA	AAG	TTI	TCG	GĢA	AAC	GAC	TTT	CTG	AAA	ACC	GGC	CAT	TAC	ACT	CAA	ATG	480
141	N	Þ	K	K	K	F	S	G	N	D	F	L	K	T	G	н	Y	T	· Q	M	160
481	GTT	TGG	GCT	'AAC	ACC	AAG	GAA	GTT	GGT	TGT	GGA	AGT.	ATA	AAA	TAC	ATT	CAA	GAG	AAA	TGG	540
161	v	W	A	N	T	ĸ	E	v	ď	C	G	S	I	K	Y	I	Q	E	K	W	180
	-	•••		•	_																
541	CAC	ααα	САТ	TAC	ירירי	GTA	TGT.	AAT	тат	GGA	CCC	AGC	GGA.	AAC	TTT	AAG	AAT	GAG	GAA	CTT	600
181	Н	ĸ	Н	Y	Ī,	v	C	N	Y	G	P	ŝ	Ĝ	N,	F	ĸ	N	E	E	L	200
-01	••	•	••	-	_	•	-		_	-	-	-	-								
601	TAT	CAA	ACA	AAG	AAT																612
	v																				204







15 B

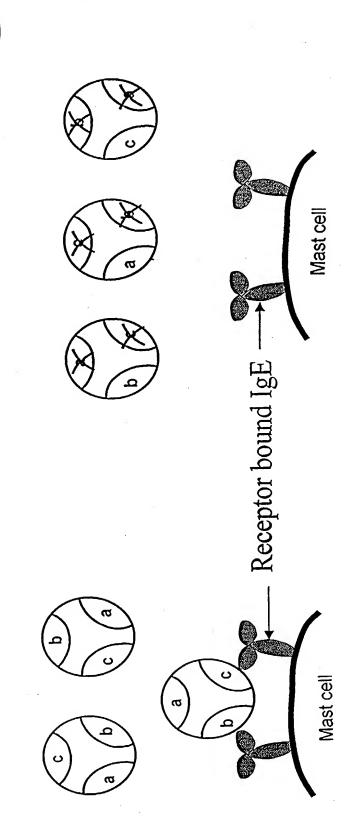




FIG. 16 A

DNA SEQUENCE

ORIGIN

cacaaattct tctttcttcc ttactactga tcattaatct gaaaacaaaa ccaaacaaac
cattcaaaat gatgtacaaa attttgtgtc tttcattgtt ggtcgcagcc gttgctcgtg
atcaagtcga tgtcaaagat tgtgccaatc atgaaatcaa aaaagttttg gtaccaggat
gccatggttc agaaccatgt atcattcatc gtggtaaacc attccaattg gaagccgttt
tcgaagccaa ccaaaacaca aaaacggcta aaattgaaat caaagcctca atcgatggtt
tagaagttga tgttcccggt atcgatccaa atgcatgcca ttacatgaaa tgcccattgg
ttaaaggaca acaatatgat attaaatata catggaatgt tccgaaaatt gcaccaaaat
ctgaaaatgt tgtcgtcact gttaaagtta tgggtgatga tggtgttttg gcctgtgcta
ttgctactca tgctaaaatc cgcgattaaa tcaaacaaaa tttattgatt ttgtaatcac
aaatgattga ttttctttcc aaaaaaaaaa taaataaaat tttgggaatt c

FIG. 16 B

mmykilclsl Ivaavardqv dvkdcanhei kkvlvpgchg sepciihrgk pfqleavfea
 nqntktakie ikasidglev dvpgidpnac hymkcplvkg qqydikytwn vpkiapksen
 vvvtvkvmgd dgvlacaiat hakird

FIG. 17

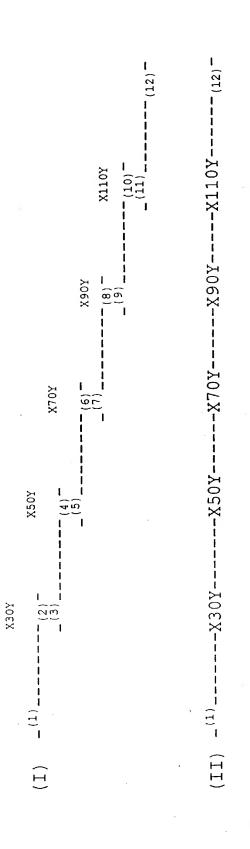






FIG. 18 A

331pMalc (s)

189BV (a)
188BV (s)

362BV (a)
361BV (s)

364BV (a)
363BV (s)

366BV (a)
365BV (s)
332pMalc (a)

DNA template: Bet v 1 (2589) carrying the Y5V mutation.

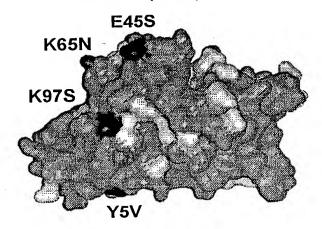
FIG. 18 B

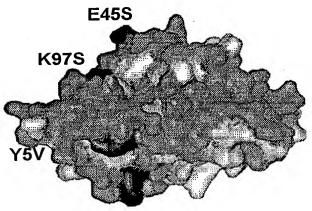
DNA template: Bet v 1 (2571) carrying N28T, K32	2Q, P108G mutations.
331pMalc	\$\$ 14.2°C
368BVa	
367BVs	•
370BVa	
<u>369BVs</u>	
	372BVa



FIG. 19 A

Bet v 1 (2628)





K134E

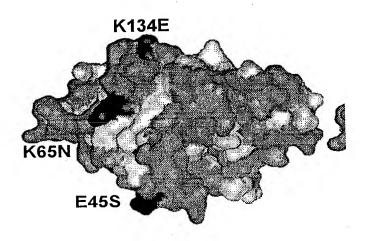
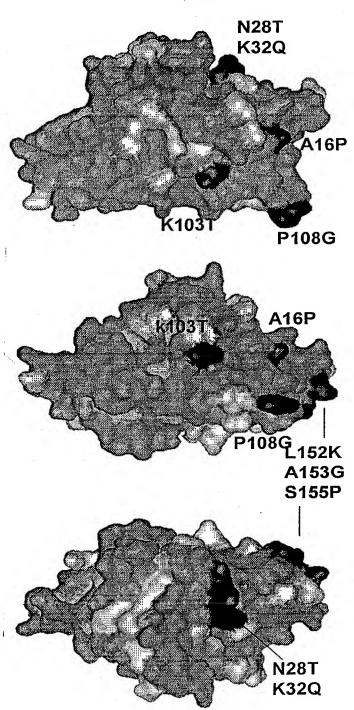


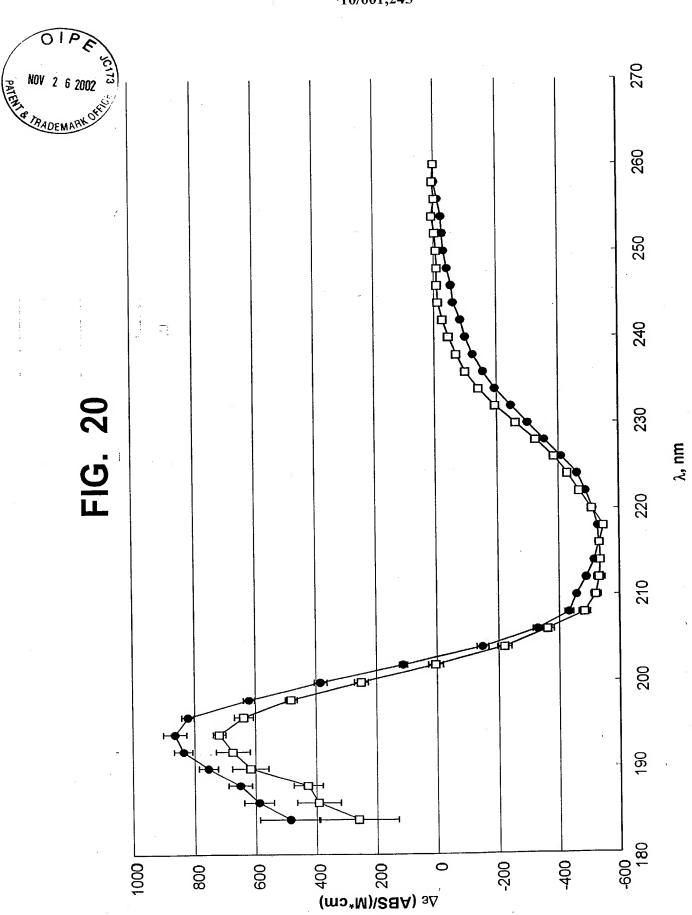


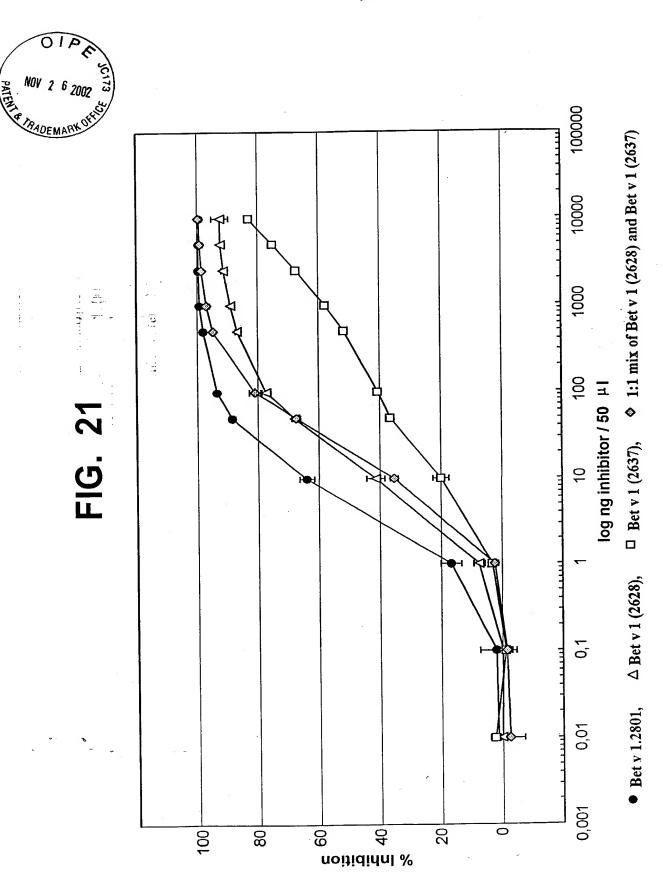
FIG. 19 B

Bet v 1 (2637)

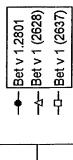


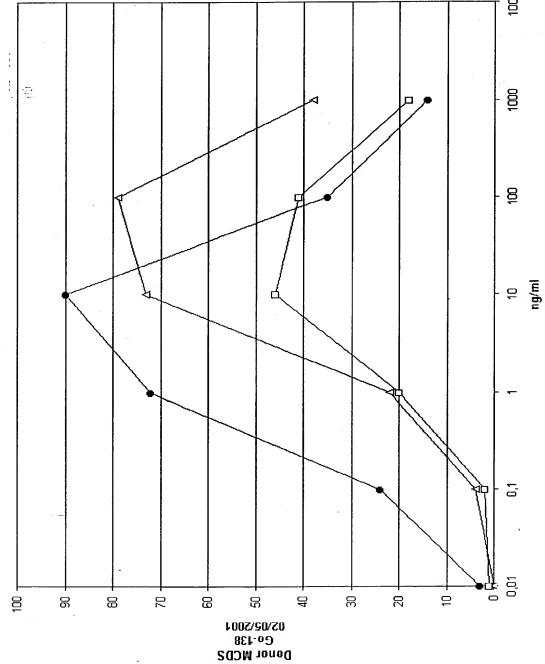




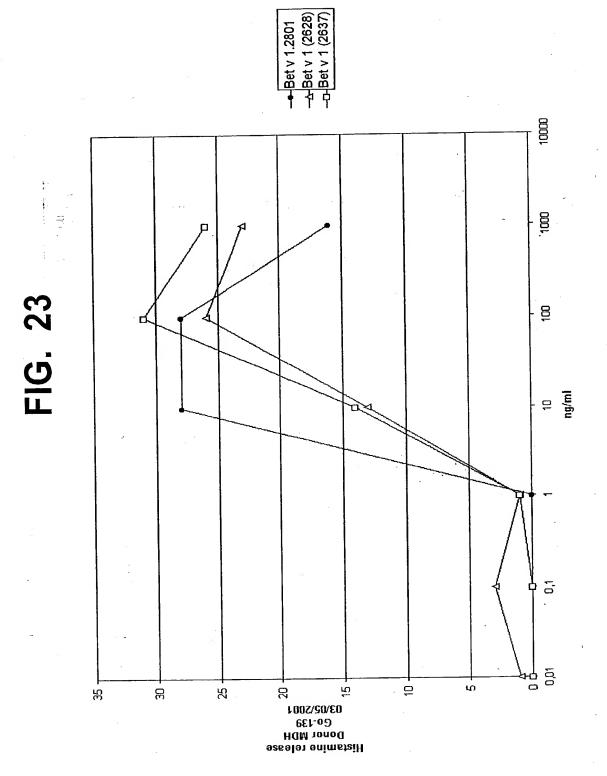


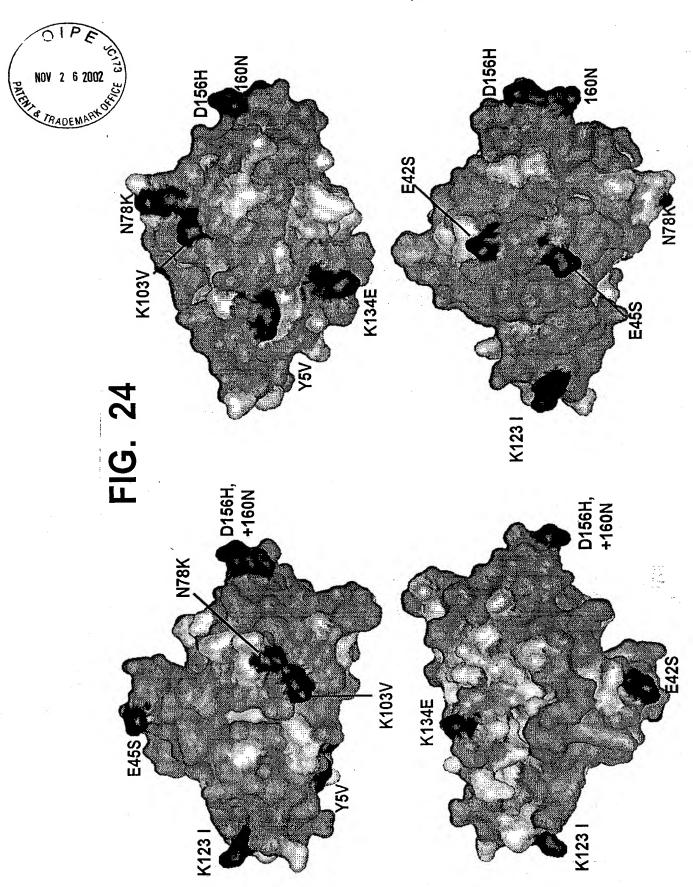


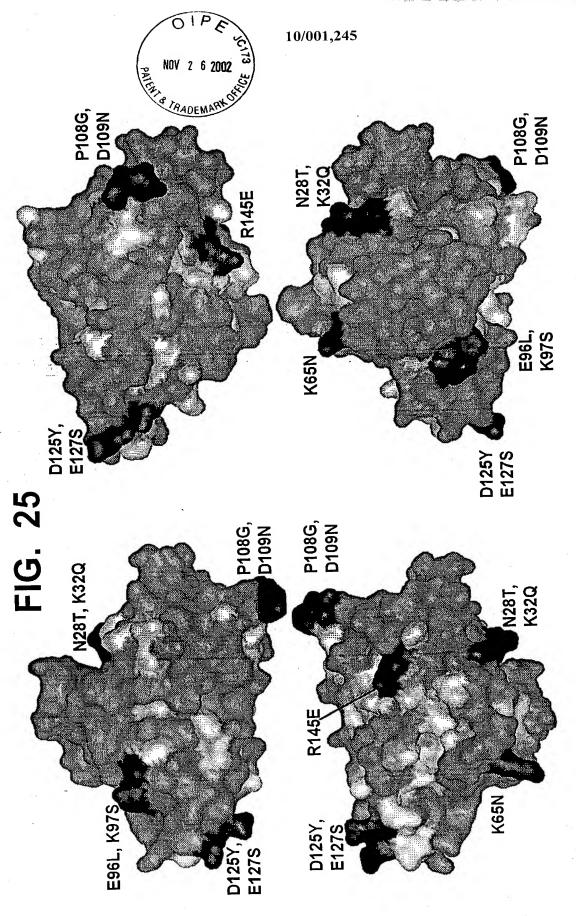


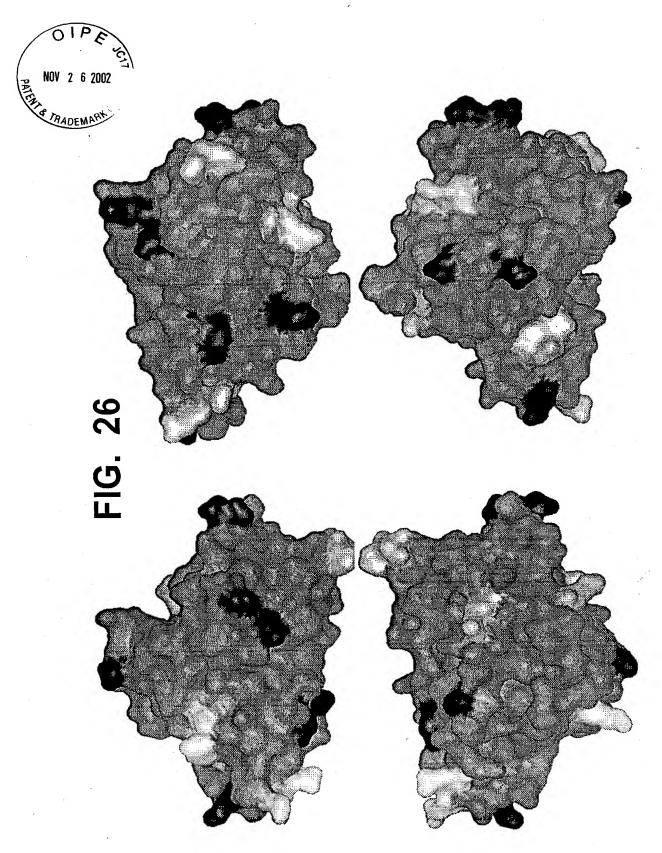


Histamine release

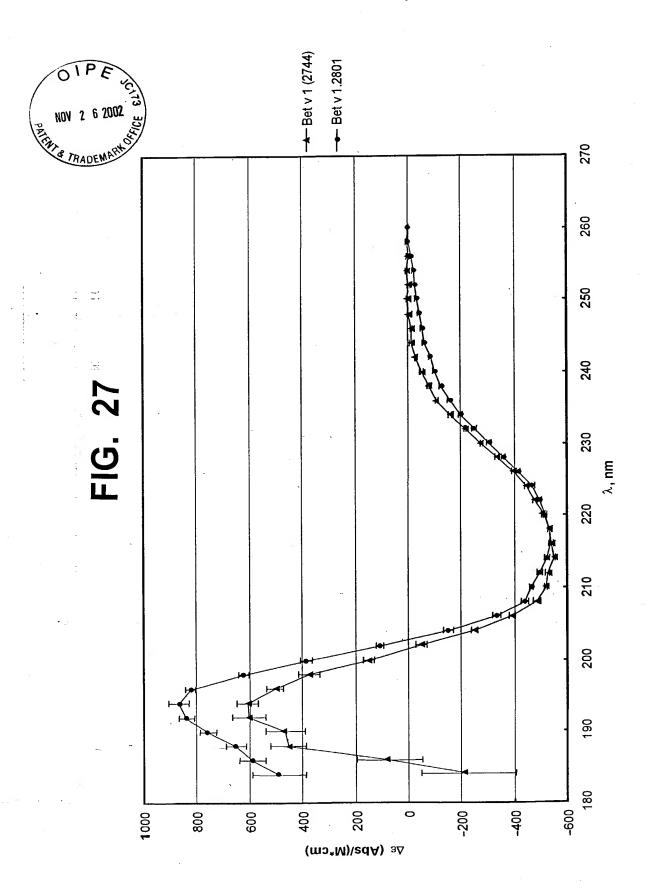


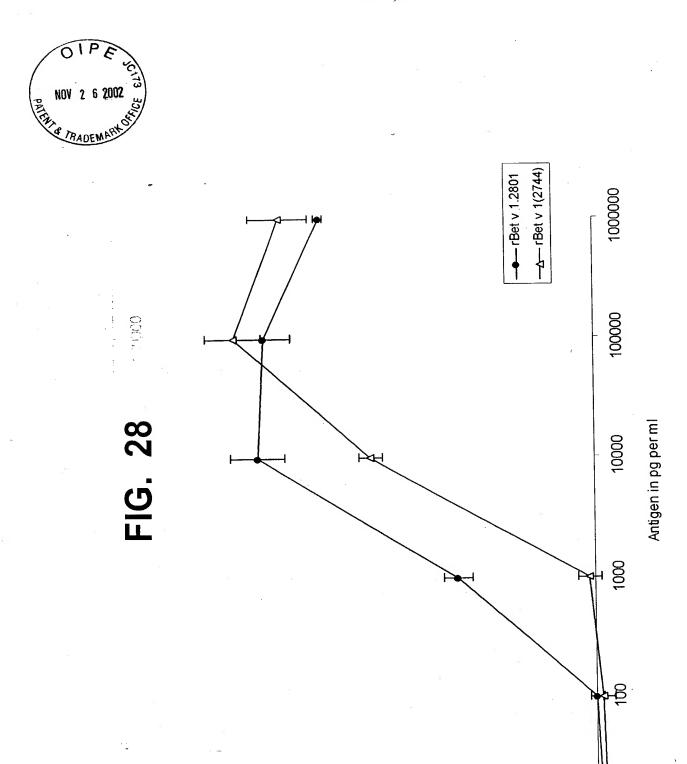






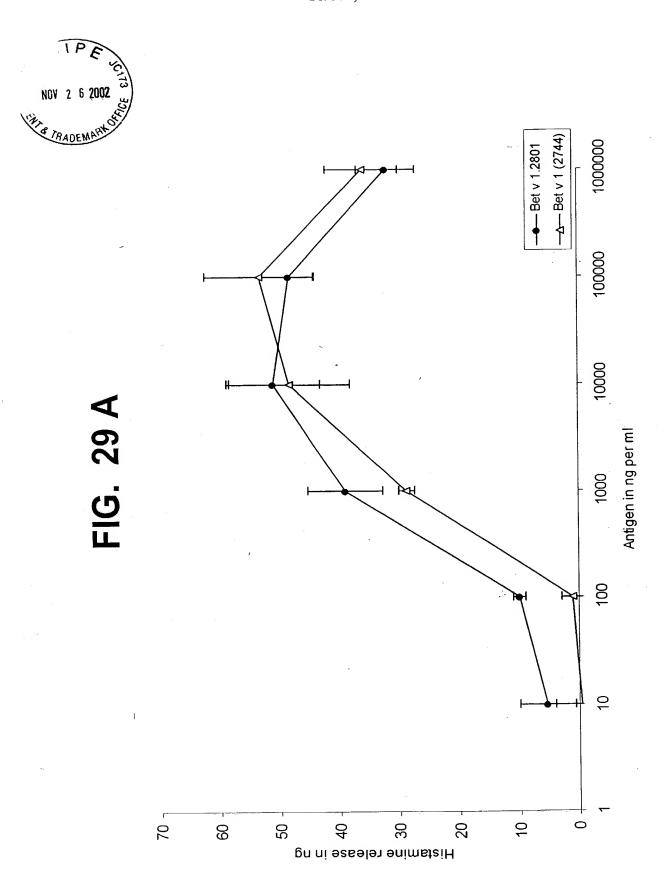
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Histamine release in ng

10/001,245



10/001,245



FIG. 29 B

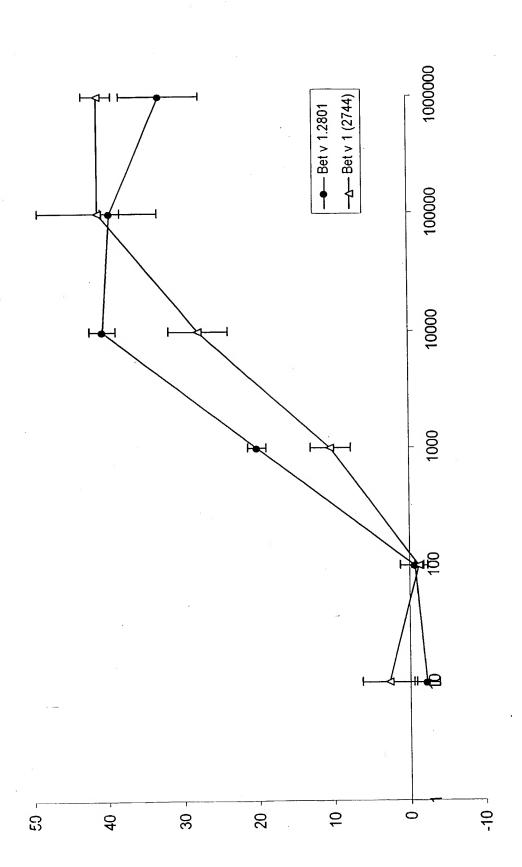
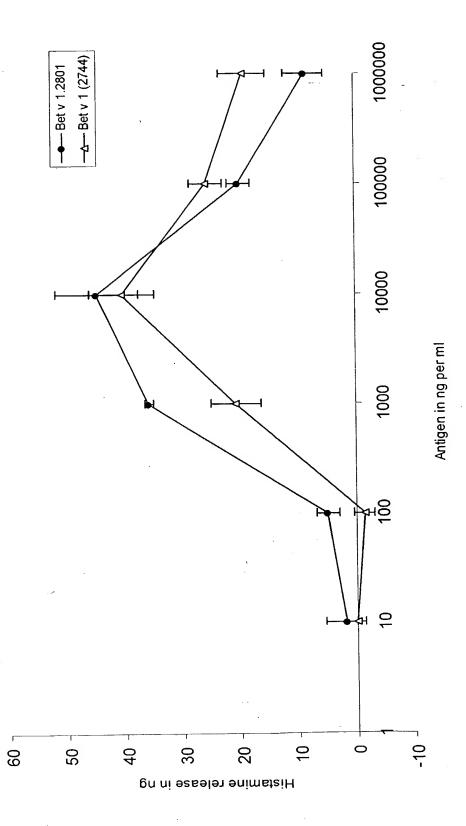




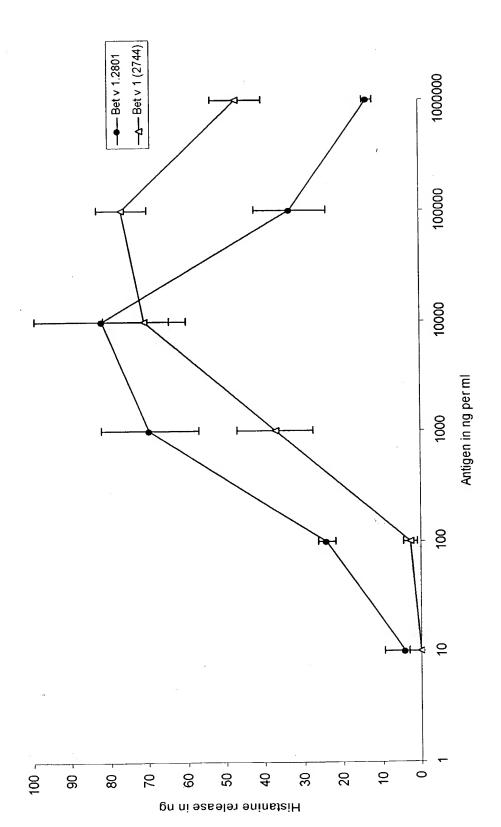
FIG. 29 C

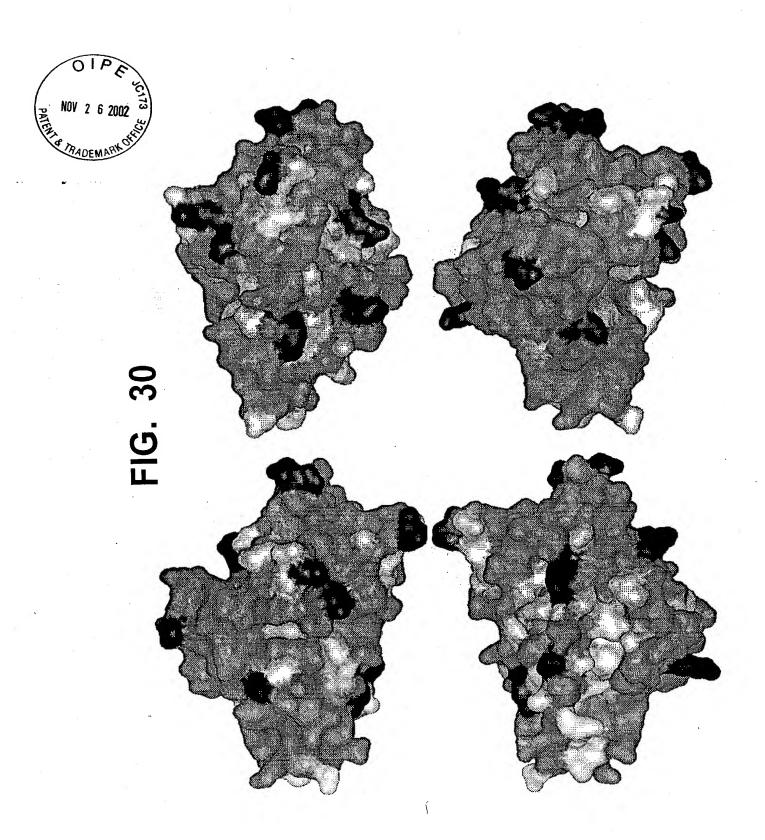


10/001,245



FIG. 29 D







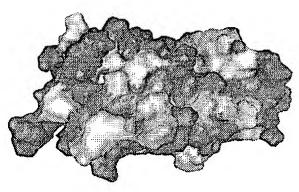
				Xho I
K6A	sense	OB43	42-тег	5' -CCGCTCGAGAAAAGAGATCAAGTCGATGTCGCCGATTGTGCC-3'
	anti-sense	OB28	39-mer	Xba I 5' -CGT <u>TCTAGA</u> CTATTAATCGCGGATTTTAGCATGAGTTGC- 3'
K15E	sense	OB44	67-mer	X 1 CCGCTCGAGAAAAGAGATCAAGTCGATGTCAAAGATTGTGCC AACCATGAAATCAAAGAAGTTTTGG-3'
	anti-sense	OB28	39-mer	Xba I 5' -CGT <u>TCTAGA</u> CTATTAATCGCGGATTTTAGCATGAGTTGC- 3'
H30N	sense	OB46	54-mer	Kpn I 5' -CGG <u>GGTACC</u> AGGATGTCATGGTTCAGAACCATGTATCATTAA CCGTGGTAAACC- 3'
	anti-sense	OB28	39-mer	3° -CGT <u>TCTAGA</u> CTATTAATCGCGGATTTTAGCATGAGTTGC-3°
E62S	sense	OB47	33-mer	5' -GCCTCAATCGATGGTTTATCAGTTGATGTTCCC-3'
	anti-sense	OB48	33-mer	5' -GGGAACATCAACTGATAAACCATCGATTGAGGC- 3'
H74N	sense	OB49	32-mer	Sph I 5' -CATG <u>GCATGC</u> AATTACATGAAATGCCCATTGG- 3'
	anti-sense	OB28	39-mer	5' -CGT <u>TCTAGA</u> CTATTAATCGCGGATTTTAGCATGAGTTGC-3'
K82N	sense	OB50	50-mer	Sph I 5'-CTACGCATGCCATTACATGAAATGCCCATTGGTTAATGGACAA CAATATG-3'
	anti-sense	OB28	39-mer	3' -CGT <u>TCTAGA</u> CTATTAATCGCGGATTTTAGCATGAGTTGC-3'

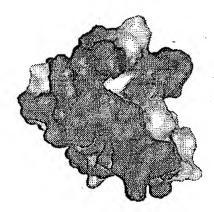
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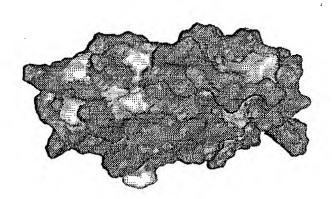
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FIG. 33







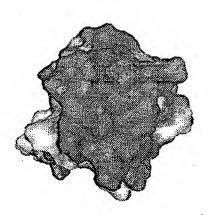
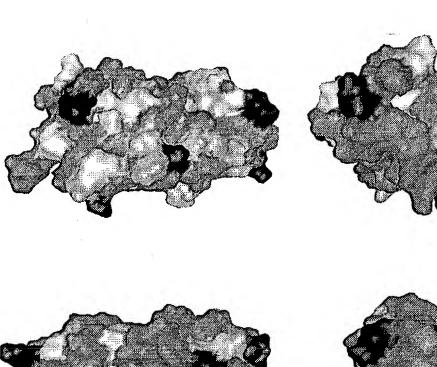
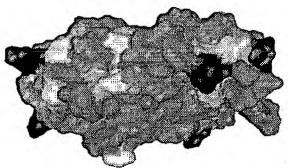
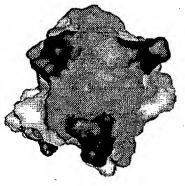




FIG. 34











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FIG. 35B

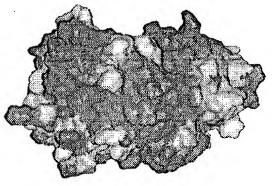
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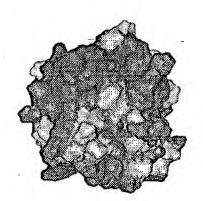
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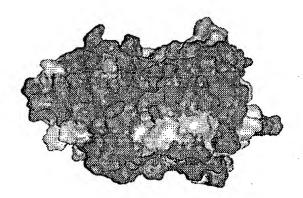
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FIG. 36







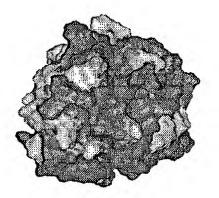
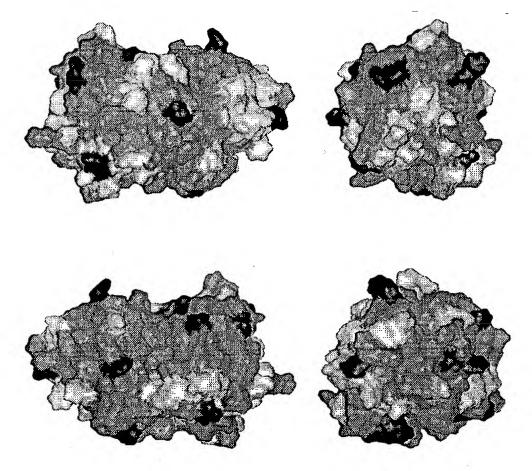




FIG. 37



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spiQ40237jMP5B_LOLPR	Poa p 5 (KBG 31) Lol p 5B Lol p 5A			ua ,	. M	A	V C	K	I N H T Y T	V	A	L I	F L	A	V	A	L V			1				•	- 1			-			.:
	Lel p 5C Phl p 5.0206				- M	A	V O	K	Y T Y T	V	A	L L	F L	A	V	A	L V L V												- -		
tr 023972 023972	Hol I 5 Phl p 5.0207		• •		- i	Α	v c	K	ΥĪ	v	A	L	F L	Ā	v	A	. V		•	 			-	-				•	:		
tn AAG42255 AAG42255 tn AAG42254 AAG42254	Hol I 58 Poa p 5		• •		- M	A	V 0		Y T	V	A	L	F L	T	v	A G	L V L V P R		•				-								
tr P93466 P93466	PM p 5.0203 PM p 5	S	V K	R	SN	G	S #		V 1	K	Δ.	A.	V P	R	R		P R P R			-				-							
spiG40963 MP5B_PHLPR triQ9SBE0 Q9SBE0	Phip 5.0204 Phip 5.02		•								-					:				-		-		- 1				-	- :		
1dO23971)O23971 sp(P56166)MP53_PHAAQ HAAQ	Pha a 5.3 Pha a 5.1	1.			- N	A	V C	ı K	Y 1 Y 1	V	A		F L	A	W V	A	L V L V			<u>.</u>			-	:			:	-,		•	
tr 004828 004828	Hor v 9 Hor v 5 (30kDa)			М	A N	S	G F	₹ E	H S	۸	٧	P	R F	₹ R	N	١.	V A	-		- i	· -		-	-							
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	PN p 5.0103				ĺ	o i	A		A 2	ۮ			2 (6 <i>)</i>			A	A P	A	G	-	-			. 6	A			6	K 7	(T	
triQ40960 Q40960 sp Q40962 MP5A_PHLPR	Phip5			-		n	Α.,	V P			¥	Ŧ.		1	P	A	A P	Â	E G	-	- -		-		A 1	0 4	À	G		X	.
8PIP22286IMP92 POAPR 8PIP22286IMP93 POAPR	Pos p 5 (KBG41)	P P	A T	L		. P	A	r (pp		S P	Α		G ۱ G ۱	4	P P	A	A P A P	A	G	-			:		A	A I	Ţ.,		k /	H	<u>.</u>
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tdP93467 P93467 sp P22284 MP91_POAPR	Phl p 5 Pos p 5 (KBG 31)	P	PΑ					K P		P #	V	G A	A \	, , ,	P	A	A F	A	Ğ T	ρ				A	A	A I	S	Ġ	K / K /	A T A T	
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tr 081344 081344 tr AAG42255 AAG42255	Phl p 5.0207 Hol I 5B		<u> </u>		•		:		-		X	A		Ţ		Â	A :	A	G	-	-		1		A	A A	A	G G	K		P P
1nJAAG42254JAAG42254 1rJOB1342JOB1342	Pos p 5 Phl p 5.0203	<u> </u>			-			: <u>-</u>	•		ľ	A A A				A	A	A	G	-			-		A		A	6	K 7 K 7	4 T 4 T	
tr/P93466 P93466 sp(040963 MP5B_PHLPR	Phi p 5 Phi p 5B										,	Α			P	Α			G	-	-		-		A	A		G	K /	A T A T	
triQ9SBE0JQ9SBE0 triOZ3971JOZ3971 spjP56166jMP53_PHAAQ	Phi p 5.0204 Phi p 5.02 Pha a 5.3								•		G	i		P 🗿		À	T		-	•	•	A	v	P		A /		Ġ	ĸ	A T	
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tr Q39995 Q39995	Hor v 5 (30kDa)					- -					- -	-	-	-							-										
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tdO81341 O81341 tdQ40960 Q40960	Phl p 5.0103 Phl p 5	E	P 1			A	A	E 8 E 5		S S	K A	Ă	Ľ	T :	K	ŀ	0 /	A A	Ϋ́Υ	K K		A \	K	Ţ			3 A	mu	P	E A	K
spiQ40962jMP5A_PHLPR spiP22285jMP92_POAPR	Phip 5 A	E	P I	(G (G		A	٨	E 6 A A	S	S S	K A N A	A V	L L	T :		L	0	4 A	Ϋ́	K	Ļ	A) A)	/ K	S	A		G A	ŧ	P	E A	K K
spiP22286[MP93_POAPR triO65319[O65319	Posp 5 (KBG60) Phlp 5	E	PI	K G K G	-1	A	Α	V D E S	S	5	K A K A	A	i.	T T	S K S K	t	D D	A A	Y	K	ì	A :		S T	A	E	G A	Ť	P	E A F A	K K
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tr 065318 065318 tr P93467 P93467	Phip5 Phip5	E	P I	K G K G K G		. A	Α	E 6 V A	S	\$ \$ \$	n A K G	A V	ì	į	5 A 5 K		E O	À ? A ?	Ý	K	ĭ	A Y	r K r K	Ť S	S A	E E	G A G A	T T	p P	E A E A	K K
spiP22284 MP91 POAPR spiQ40237 MP5B_LOLPR	Lolp 5B Lolp 5A		e te	- G			A	D .	Q	5	K N	Q	ĭ	1	\$ * P *	(<u>L</u>	D. D	A #		K	i. V	A Y	Y E Y K	A	A	F	G #	, T	P	E A F A	K
tr Q9XF24 Q9XF24 tr Q9SC99 Q9SC99 tr Q81343 Q81343	Lel p 5C Phl p 5 0206					. D	A	. A	P P	-		G G	i L		L II P II	(L	D D	T [١Y	S	٧ ٧	A .	Y # Y #	A	G A	E V	G A G A	Ţ	P	E A	K K
tr O81343 O01345 tr O81344 O81344	Hol 15 Phi p 5.0207					A	T	K A	P	- [G	*****	٧		l L	Ŋ	A I	ŧΥ	S		S	Ý	Α	A	Q V V	Ğ İ	Ĭ	P	E A	K
tn AAG42255 AAG42255 tn AAG42254 AAG42254	Hol I 5B Poa p 5			- -			A	Q A	P	-		G		٧		1 V	A A D	A I A I	S	D	A	T	Y	ĢΑ	Α	٧	G A	ij	p p	Ë ¢	K
tr 061342 061342 tr P93466 P93466	Phlp 5 0203 Phlp 5	1						. p	P			G G		٧	P I	(L	D D	À			8	Ā A		Α	Α	٧			P P	E #	, K , K
sp 040963 MP5B_PHLP6 tr 095BE0 095BE0	Phip 568 Phip 5.0204 Phip 5.02			- -			Ė	 T .	P	P		G	m	٧	Р ј 5 ј	(L	0	A D	i Y N Y	S	V	A A	Y	A	Α	٧	G /	(T	p Q	E #	K
1r O23971 O23971 sp P56166 MP53_PHAAG HAAG					-		S	T P	G	P	•	Q	18	÷	A 21	·	D	A	m	S R	V	V A	Y 1	Α	A	A E	G G	Ţ	P	£ £	K
tr 004828 004828 - tr 039995 039995	Hor v 9 Hor v 5 (30kDa)		\ A	T 🥳	۵	!	Ğ	Q			M	K	il.	S	S S	S (L	E	L :	S S	K	ľ	Å	Y C) K	Â	0	Ğ	ì	P	E ;	K

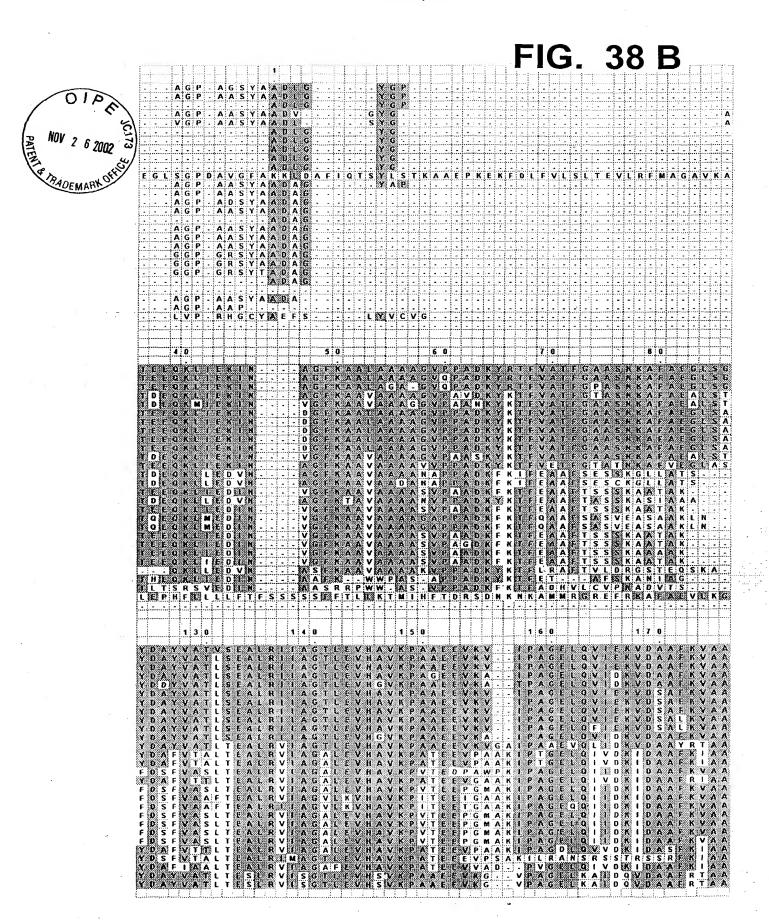




FIG. 38 C

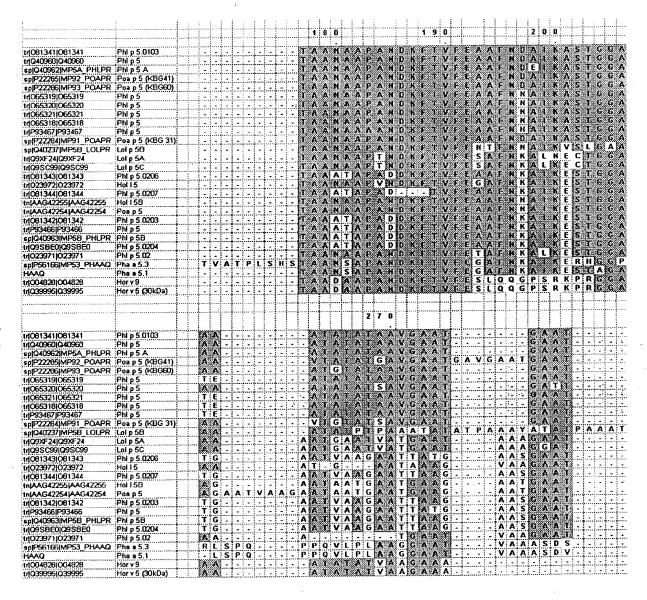




FIG. 38 D

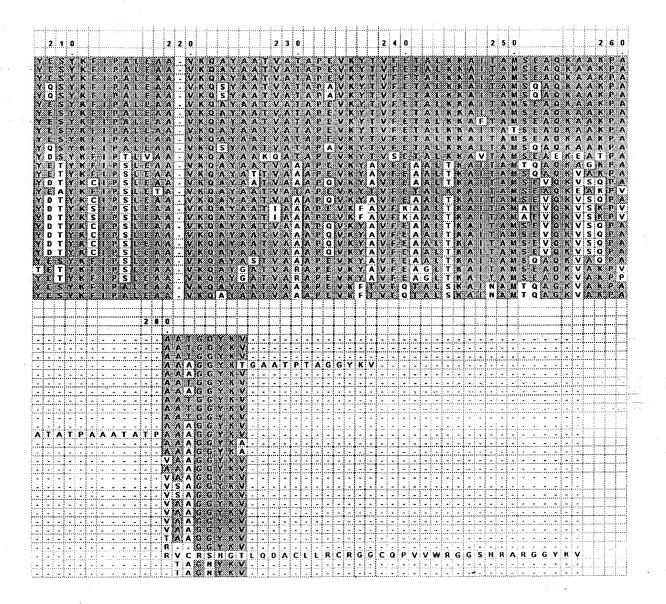
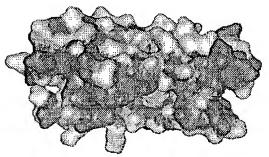
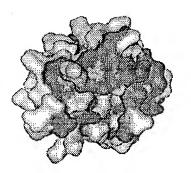
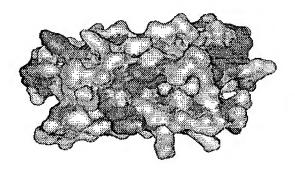




FIG. 39 A







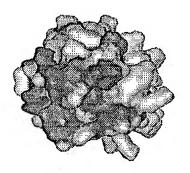
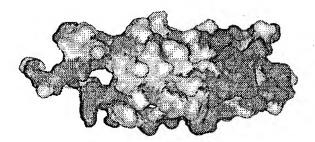
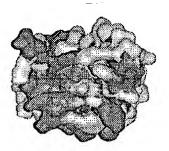
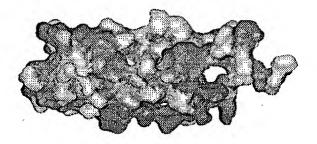




FIG. 39 B







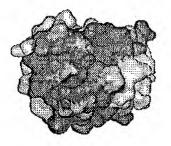
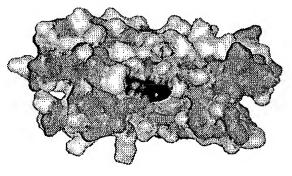
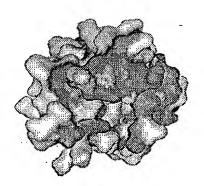
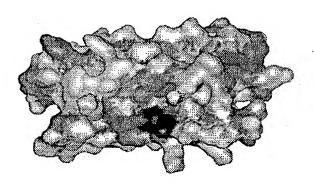




FIG. 40 A







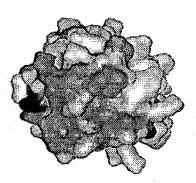
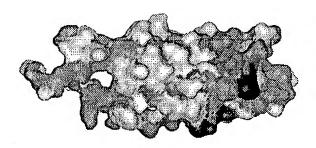
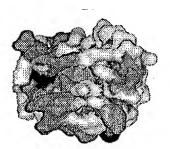
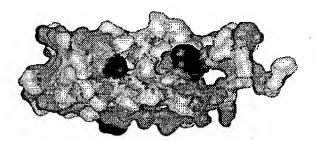


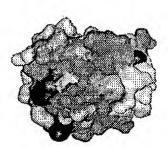


FIG. 40 B

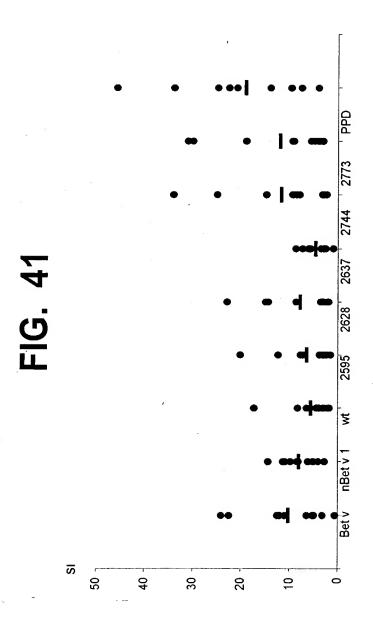






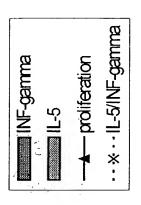






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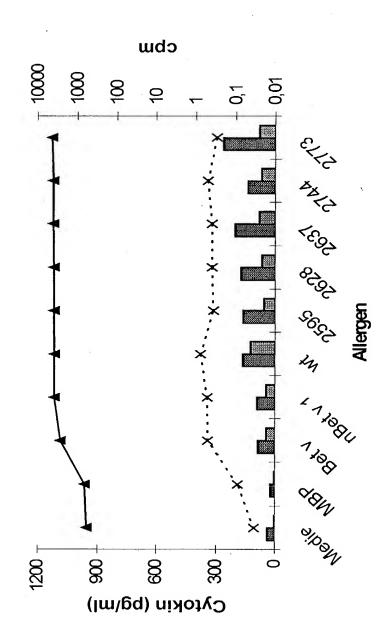


FIG. 42

